

1642  
PH#12



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See page 6



1600

RAW SEQUENCE LISTING

DATE: 06/03/2002

PATENT APPLICATION: US/09/492,764B

TIME: 13:31:22

Input Set : A:\10873-008.txt

Output Set: N:\CRF3\06032002\I492764B.raw

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JUN 12 2002

TECH CENTER 1600/2900

```

5 <110> APPLICANT: Jove, Richard
7   Hamilton, Andrew
9   Gilbert, Richard
11  Dalton, William
13  Sebti, Said
15  Yu, Hua
17  Heller, Richard
19  Jaroszeski, Mark
23 <120> TITLE OF INVENTION: INHIBITION OF STAT3 SIGNAL TRANSDUCTION FOR HUMAN CANCER
24  THERAPY
28 <130> FILE REFERENCE: 10873-008-999
32 <140> CURRENT APPLICATION NUMBER: 09/492,764B
34 <141> CURRENT FILING DATE: 2000-01-27
38 <150> PRIOR APPLICATION NUMBER: 60/117,600
40 <151> PRIOR FILING DATE: 1999-01-27
44 <160> NUMBER OF SEQ ID NOS: 38
48 <170> SOFTWARE: PatentIn version 3.0
52 <210> SEQ ID NO: 1
54 <211> LENGTH: 24
56 <212> TYPE: DNA
58 <213> ORGANISM: Homo sapiens
62 <400> SEQUENCE: 1
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66 <210> SEQ ID NO: 2
68 <211> LENGTH: 24
70 <212> TYPE: DNA
72 <213> ORGANISM: Homo sapiens
76 <400> SEQUENCE: 2
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80 <210> SEQ ID NO: 3
82 <211> LENGTH: 22
84 <212> TYPE: DNA
86 <213> ORGANISM: Homo sapiens
90 <400> SEQUENCE: 3
91 gtcccccggc cggggaggcg ct                               22
94 <210> SEQ ID NO: 4
96 <211> LENGTH: 25
98 <212> TYPE: DNA
100 <213> ORGANISM: Homo sapiens
104 <400> SEQUENCE: 4
105 cgacgacttc tcccgccgct accgc                           25
108 <210> SEQ ID NO: 5
110 <211> LENGTH: 25

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 Output Set: N:\CRF3\06032002\I492764B.raw

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112 <212> TYPE: DNA
114 <213> ORGANISM: Homo sapiens
118 <400> SEQUENCE: 5
119 ccgcatgctg gggccgtaca gttcc 25
122 <210> SEQ ID NO: 6
124 <211> LENGTH: 20
126 <212> TYPE: DNA
128 <213> ORGANISM: Homo sapiens
132 <400> SEQUENCE: 6
133 cgggcattca gtgacctgac 20
136 <210> SEQ ID NO: 7
138 <211> LENGTH: 20
140 <212> TYPE: DNA
142 <213> ORGANISM: Homo sapiens
146 <400> SEQUENCE: 7
147 tcaggaacca gcggttgaag 20
150 <210> SEQ ID NO: 8
152 <211> LENGTH: 20
154 <212> TYPE: DNA
156 <213> ORGANISM: Homo sapiens
160 <400> SEQUENCE: 8
161 ccactgaact tctgattcgc 20
164 <210> SEQ ID NO: 9
166 <211> LENGTH: 20
168 <212> TYPE: DNA
170 <213> ORGANISM: Homo sapiens
174 <400> SEQUENCE: 9
175 gcgtgctagc tggatgtctt 20
178 <210> SEQ ID NO: 10
180 <211> LENGTH: 9
182 <212> TYPE: DNA
184 <213> ORGANISM: Homo sapiens
188 <400> SEQUENCE: 10
189 ttcggagaa 9
192 <210> SEQ ID NO: 11
194 <211> LENGTH: 9
196 <212> TYPE: DNA
198 <213> ORGANISM: Homo sapiens
202 <400> SEQUENCE: 11
203 tgaggataa 9
206 <210> SEQ ID NO: 12
208 <211> LENGTH: 12
210 <212> TYPE: PRT
212 <213> ORGANISM: Homo sapiens
216 <220> FEATURE:
218 <221> NAME/KEY: misc_feature
220 <223> OTHER INFORMATION: Xaa = Ser or Pro
224 <400> SEQUENCE: 12
W--> 226 His Tyr Xaa Pro Ile Leu Val Tyr Gln Pro Ser Trp

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227 1          5          10
229 <210> SEQ ID NO: 13
231 <211> LENGTH: 12
233 <212> TYPE: PRT
235 <213> ORGANISM: Homo sapiens
239 <400> SEQUENCE: 13
241 Gln Asp Val His Leu Thr Gln Gln Ser Arg Tyr Thr
242 1          5          10
244 <210> SEQ ID NO: 14
246 <211> LENGTH: 12
248 <212> TYPE: PRT
250 <213> ORGANISM: Homo sapiens
254 <400> SEQUENCE: 14
256 Ser His Pro Trp Asn Ala Gln Arg Glu Leu Ser Val
257 1          5          10
259 <210> SEQ ID NO: 15
261 <211> LENGTH: 12
263 <212> TYPE: PRT
265 <213> ORGANISM: Homo sapiens
269 <400> SEQUENCE: 15
271 Tyr Pro Ala Pro Gln Pro Leu Val Thr Lys Thr Ser
272 1          5          10
274 <210> SEQ ID NO: 16
276 <211> LENGTH: 12
278 <212> TYPE: PRT
280 <213> ORGANISM: Homo sapiens
284 <400> SEQUENCE: 16
286 Phe Ser Tyr Pro Leu Thr Arg Ala Pro Leu Asn Met
287 1          5          10
289 <210> SEQ ID NO: 17
291 <211> LENGTH: 7
293 <212> TYPE: PRT
295 <213> ORGANISM: Homo sapiens
299 <400> SEQUENCE: 17
301 His Ala Ile Tyr Pro Arg Asn
302 1          5
304 <210> SEQ ID NO: 18
306 <211> LENGTH: 7
308 <212> TYPE: PRT
310 <213> ORGANISM: Homo sapiens
314 <400> SEQUENCE: 18
316 Ala Ser Thr Leu Pro Lys Ala
317 1          5
319 <210> SEQ ID NO: 19
321 <211> LENGTH: 7
323 <212> TYPE: PRT
325 <213> ORGANISM: Homo sapiens
329 <400> SEQUENCE: 19
331 Ile Gln Ser Pro His Phe Phe
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## RAW SEQUENCE LISTING

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Input Set : A:\10873-008.txt

Output Set: N:\CRF3\06032002\I492764B.raw

```
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334 <210> SEQ ID NO: 20
336 <211> LENGTH: 6
338 <212> TYPE: PRT
340 <213> ORGANISM: Homo sapiens
344 <220> FEATURE:
346 <221> NAME/KEY: misc_feature
348 <223> OTHER INFORMATION: X = PHOSPHOTYROSINE
352 <400> SEQUENCE: 20
W--> 354 Pro Xaa Leu Lys Thr Lys
355 1          5
357 <210> SEQ ID NO: 21
359 <211> LENGTH: 6
361 <212> TYPE: PRT
363 <213> ORGANISM: Homo sapiens
367 <400> SEQUENCE: 21
369 Pro Tyr Leu Lys Thr Lys
370 1          5
372 <210> SEQ ID NO: 22
374 <211> LENGTH: 6
376 <212> TYPE: PRT
378 <213> ORGANISM: Homo sapiens
382 <220> FEATURE:
384 <221> NAME/KEY: misc_feature
386 <223> OTHER INFORMATION: X= PHOSPHOTYROSINE
390 <400> SEQUENCE: 22
W--> 392 Ala Xaa Leu Lys Thr Lys
393 1          5
395 <210> SEQ ID NO: 23
397 <211> LENGTH: 6
399 <212> TYPE: PRT
401 <213> ORGANISM: Homo sapiens
405 <220> FEATURE:
407 <221> NAME/KEY: misc_feature
409 <223> OTHER INFORMATION: X = PHOSPHOTYROSINE
413 <400> SEQUENCE: 23
W--> 415 Pro Xaa Ala Lys Thr Lys
416 1          5
418 <210> SEQ ID NO: 24
420 <211> LENGTH: 6
422 <212> TYPE: PRT
424 <213> ORGANISM: Homo sapiens
428 <220> FEATURE:
430 <221> NAME/KEY: misc_feature
432 <223> OTHER INFORMATION: X = PHOSPHOTYROSINE
436 <400> SEQUENCE: 24
W--> 438 Pro Xaa Leu Ala Thr Lys
439 1          5
441 <210> SEQ ID NO: 25
```

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/492,764B

DATE: 06/03/2002  
TIME: 13:31:22

Input Set : A:\10873-008.txt  
Output Set: N:\CRF3\06032002\I492764B.raw

443 <211> LENGTH: 6  
445 <212> TYPE: PRT  
447 <213> ORGANISM: Homo sapiens  
451 <220> FEATURE:  
453 <221> NAME/KEY: misc\_feature  
455 <223> OTHER INFORMATION: X = PHOSPHOTYROSINE  
459 <400> SEQUENCE: 25  
W--> 461 Pro Xaa Leu Lys Ala Lys  
462 1 5  
464 <210> SEQ ID NO: 26  
466 <211> LENGTH: 6  
468 <212> TYPE: PRT  
470 <213> ORGANISM: Homo sapiens  
474 <400> SEQUENCE: 26  
476 Pro Tyr Leu Lys Thr Ala  
477 1 5  
479 <210> SEQ ID NO: 27  
481 <211> LENGTH: 4  
483 <212> TYPE: PRT  
485 <213> ORGANISM: Homo sapiens  
489 <220> FEATURE:  
491 <221> NAME/KEY: misc\_feature  
493 <223> OTHER INFORMATION: X = PHOSPHOTYROSINE  
497 <400> SEQUENCE: 27  
W--> 499 Pro Xaa Leu Lys  
500 1  
502 <210> SEQ ID NO: 28  
504 <211> LENGTH: 4  
506 <212> TYPE: PRT  
508 <213> ORGANISM: Homo sapiens  
512 <220> FEATURE:  
514 <221> NAME/KEY: misc\_feature  
516 <223> OTHER INFORMATION: X = PHOSPHOTYROSINE  
520 <400> SEQUENCE: 28  
W--> 522 Pro Xaa Phe Lys  
523 1  
525 <210> SEQ ID NO: 29  
527 <211> LENGTH: 3  
529 <212> TYPE: PRT  
531 <213> ORGANISM: Homo sapiens  
535 <220> FEATURE:  
537 <221> NAME/KEY: misc\_feature  
539 <223> OTHER INFORMATION: X = PHOSPHOTYROSINE  
543 <400> SEQUENCE: 29  
W--> 545 Xaa Leu Lys  
546 1  
548 <210> SEQ ID NO: 30  
550 <211> LENGTH: 4  
552 <212> TYPE: PRT

RAW SEQUENCE LISTING ERROR SUMMARY      DATE: 06/03/2002  
PATENT APPLICATION: US/09/492,764B      TIME: 13:31:23

Input Set : A:\10873-008.txt  
Output Set: N:\CRF3\06032002\I492764B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:12; Xaa Pos. 3  
Seq#:20; Xaa Pos. 2  
Seq#:22; Xaa Pos. 2  
Seq#:23; Xaa Pos. 2  
Seq#:24; Xaa Pos. 2  
Seq#:25; Xaa Pos. 2  
Seq#:27; Xaa Pos. 2  
Seq#:28; Xaa Pos. 2  
Seq#:29; Xaa Pos. 1  
Seq#:30; Xaa Pos. 2  
Seq#:32; Xaa Pos. 2  
Seq#:34; Xaa Pos. 2  
Seq#:35; Xaa Pos. 2  
Seq#:36; Xaa Pos. 2  
Seq#:37; Xaa Pos. 2  
Seq#:38; Xaa Pos. 2

VARIABLE LOCATION SUMMARY  
PATENT APPLICATION: US/09/492,764B

DATE: 06/03/2002  
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Input Set : A:\10873-008.txt  
Output Set: N:\CRF3\06032002\I492764B.raw

Use of n's or Xaa's(NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.  
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.  
in <220> to <223> section, please explain location of n or Xaa, and which  
residue n or Xaa represents.

Seq#:12; Xaa Pos. 3  
Seq#:20; Xaa Pos. 2  
Seq#:22; Xaa Pos. 2  
Seq#:23; Xaa Pos. 2  
Seq#:24; Xaa Pos. 2  
Seq#:25; Xaa Pos. 2  
Seq#:27; Xaa Pos. 2  
Seq#:28; Xaa Pos. 2  
Seq#:29; Xaa Pos. 1  
Seq#:30; Xaa Pos. 2  
Seq#:32; Xaa Pos. 2  
Seq#:34; Xaa Pos. 2  
Seq#:35; Xaa Pos. 2  
Seq#:36; Xaa Pos. 2  
Seq#:37; Xaa Pos. 2  
Seq#:38; Xaa Pos. 2

## VERIFICATION SUMMARY

DATE: 06/03/2002

PATENT APPLICATION: US/09/492,764B

TIME: 13:31:23

Input Set : A:\10873-008.txt

Output Set: N:\CRF3\06032002\I492764B.raw

L:226 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:12  
L:226 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0  
L:354 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:20  
L:354 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0  
L:392 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:22  
L:392 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0  
L:415 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:23  
L:415 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0  
L:438 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:24  
L:438 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0  
L:461 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:25  
L:461 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0  
L:499 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:27  
L:499 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0  
L:522 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:28  
L:522 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0  
L:545 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:29  
L:545 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:0  
L:568 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:30  
L:568 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0  
L:606 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:32  
L:606 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0  
L:644 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:34  
L:644 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:0  
L:667 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:35  
L:667 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:0  
L:690 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:36  
L:690 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:0  
L:713 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:37  
L:713 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:0  
L:736 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:38  
L:736 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:0